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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/940,925A

DATE: 06/21/2002
TIME: 12:20:02

Input Set : A:\Seqsub2.app
Output Set: N:\CRF3\06212002\I940925A.raw

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       SEQUENCE LISTING
 4 (1) GENERAL INFORMATION:
 6      (i) APPLICANT: BROW, MARY ANN D.
 7                     LYAMICHEV, VICTOR I.
 8                     OLIVE, DAVID M.
10     (ii) TITLE OF INVENTION: RAPID DETECTION AND IDENTIFICATION OF
11                     PATHOGENS
13    (iii) NUMBER OF SEQUENCES: 165
15     (iv) CORRESPONDENCE ADDRESS:
16          (A) ADDRESSEE: MEDLEN & CARROLL
17          (B) STREET: 220 MONTGOMERY STREET, SUITE 2200
18          (C) CITY: SAN FRANCISCO
19          (D) STATE: CALIFORNIA
20          (E) COUNTRY: UNITED STATES OF AMERICA
21          (F) ZIP: 94104
23      (v) COMPUTER READABLE FORM:
24          (A) MEDIUM TYPE: Floppy disk
25          (B) COMPUTER: IBM PC compatible
26          (C) OPERATING SYSTEM: PC-DOS/MS-DOS
27          (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
29     (vi) CURRENT APPLICATION DATA:
30          (A) APPLICATION NUMBER: US/09/940,925A
31          (B) FILING DATE: 10-Jun-2002
32          (C) CLASSIFICATION:
34   (viii) ATTORNEY/AGENT INFORMATION:
35          (A) NAME: CARROLL, PETER G.
36          (B) REGISTRATION NUMBER: 32,837
37          (C) REFERENCE/DOCKET NUMBER: FORS-01756
39     (ix) TELECOMMUNICATION INFORMATION:
40          (A) TELEPHONE: (415) 705-8410
41          (B) TELEFAX: (415) 397-8338
44 (2) INFORMATION FOR SEQ ID NO: 1:
46      (i) SEQUENCE CHARACTERISTICS:
47          (A) LENGTH: 2506 base pairs
48          (B) TYPE: nucleic acid
49          (C) STRANDEDNESS: double
50          (D) TOPOLOGY: linear
52     (ii) MOLECULE TYPE: DNA (genomic)
56     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
58 ATGAGGGGGA TGCTGCCCCT CTTTGAGCCC AAGGGCCGGG TCCTCCTGGT GGACGGCCAC        60
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62 GTGCAGGCGG TCTACGGCTT CGCCAAGAGC CTCCTCAAGG CCCTCAAGGA GGACGGGGAC       180
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C--> 30

C--> 31

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144      (i) SEQUENCE CHARACTERISTICS:
145          (A) LENGTH: 2496 base pairs
146          (B) TYPE: nucleic acid
147          (C) STRANDEDNESS: double
148          (D) TOPOLOGY: linear
150     (ii) MOLECULE TYPE: DNA (genomic)
154     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
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160 CAGGCGGTCT ACGGCTTCGC CAAAAGCCTC CTCAAGGCCC TGAAGGAGGA CGGGGACGTG    180
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240 (2) INFORMATION FOR SEQ ID NO: 3:
242       (i) SEQUENCE CHARACTERISTICS:
243             (A) LENGTH: 2504 base pairs
244             (B) TYPE: nucleic acid
245             (C) STRANDEDNESS: double
246             (D) TOPOLOGY: linear
248       (ii) MOLECULE TYPE: DNA (genomic)
252       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
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324 GCTTCCCCAA GGTGCGGGCC TGGATAGAAA AGACCCTGGA GGAGGGGAGG AAGCGGGGCT    2160
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330 ACCTCATGAA GCTCGCCATG GTGAAGCTCT TCCCCCGCCT CCGGGAGATG GGGGCCCGCA    2340
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334 TGGCGGCTTT GGCCAAGGAG GCCATGGAGA AGGCCTATCC CCTCGCCGTG CCCCTGGAGG    2460
336 TGGAGGTGGG GATGGGGGAG GACTGGCTTT CCGCCAAGGG TTAG               2504
338 (2) INFORMATION FOR SEQ ID NO: 4:
340     (i) SEQUENCE CHARACTERISTICS:
341         (A) LENGTH: 832 amino acids
342         (B) TYPE: amino acid
343         (C) STRANDEDNESS: single
344         (D) TOPOLOGY: linear
346     (ii) MOLECULE TYPE: protein
350     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
352     Met Arg Gly Met Leu Pro Leu Phe Glu Pro Lys Gly Arg Val Leu Leu
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365    65                  70              .       75                 80
367    Tyr Lys Ala Gly Arg Ala Pro Thr Pro Glu Asp Phe Pro Arg Gln Leu
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371                  100                 105                110
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382    Tyr Leu Ile Thr Pro Ala Trp Leu Trp Glu Lys Tyr Gly Leu Arg Pro
383                 .165                 170                175
385    Asp Gln Trp Ala Asp Tyr Arg Ala Leu Thr Gly Asp Glu Ser Asp Asn
386              180                 185                190
388    Leu Pro Gly Val Lys Gly Ile Gly Glu Lys Thr Ala Arg Lys Leu Leu
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391    Glu Glu Trp Gly Ser Leu Glu Ala Leu Leu Lys Asn Leu Asp Arg Leu
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395    225                 230                235                240
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401              260                 265                270
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413              325                 330                335
415    Glu Pro Tyr Lys Ala Leu Arg Asp Leu Lys Glu Ala Arg Gly Leu Leu
416              340                 345                350
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